

SEQUENCE LISTING

<110> Blaschuk, Orest W.
Michaud, Stephanie D.

<120> COMPOUNDS AND METHODS FOR MODULATING
FUNCTIONS OF NONCLASSICAL CADHERINS

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Met Trp Asn Gln Phe Phe
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cadherin Trp-containing CAR sequence

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<223> Xaa = Phe, Ala or Ile

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<223> Xaa = Ala or Pro

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Xaa Trp Xaa Xaa Xaa Xaa
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<223> Representative desmosomal cadherin Trp-containing
CAR sequence

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CAR sequence

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CAR sequence

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<210> 1296

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1296

Glu Ala Trp Ile Thr Ala Pro Lys

1

5

<210> 1297

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1297

Glu Ala Trp Ile Thr Ala Pro Val Lys

1

5

<210> 1298

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1298

Glu Ala Trp Ile Thr Ala Pro Val Ala Lys
1 5 10

<210> 1299

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1299

Glu Ala Trp Ile Thr Ala Pro Val Ala Leu Lys
1 5 10

<210> 1300

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1300

Glu Trp Ile Thr Lys
1 5

<210> 1301

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1301

Glu Trp Ile Thr Ala Lys
1 5

<210> 1302

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1302

Glu Trp Ile Thr Ala Pro Lys

1 5

<210> 1303

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1303

Glu Trp Ile Thr Ala Pro Val Lys

1 5

<210> 1304

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1304

Glu Trp Ile Thr Ala Pro Val Ala Lys

1 5

<210> 1305

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Exemplary cyclic peptide

<400> 1305

Glu Trp Ile Thr Ala Pro Val Ala Leu Lys

1 5 10

<210> 1306

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Preferred CAR sequence for inclusion with a
modulating agent

<400> 1306
 Tyr Ile Gly Ser Arg
 1 5

<210> 1307
 <211> 10
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Preferred CAR sequence for inclusion with a
 modulating agent

<400> 1307
 Lys Tyr Ser Phe Asn Tyr Asp Gly Ser Glu
 1 5 10

<210> 1308
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Preferred CAR sequence for inclusion with a
 modulating agent

<400> 1308
 Ser Phe Thr Ile Asp Pro Lys Ser Gly
 1 5

<210> 1309
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Preferred CAR sequence for inclusion with a
 modulating agent

<400> 1309
 Leu Tyr His Tyr
 1

<210> 1310
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Claudin CAR sequence comprising at least four

consecutive amino acids present within a claudin region

<221> VARIANT

<222> 2

<223> Xaa = Lys or Arg

<221> VARIANT

<222> 3

<223> Xaa = any amino acid

<221> VARIANT

<222> 4

<223> Xaa = any amino acid

<221> VARIANT

<222> 5

<223> Xaa = Ser or Ala

<221> VARIANT

<222> 6

<223> Xaa = Tyr or Phe

<221> VARIANT

<222> (7)...(7)

<223> Xaa = any amino acid

<400> 1310

Trp Xaa Xaa Xaa Xaa Xaa Xaa Gly
1 5

<210> 1311

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Atypical cadherin CAR sequence comprising at least three consecutive amino acids present within an atypical cadherin region

<221> VARIANT

<222> 1,3

<223> Xaa = any amino acid

<221> VARIANT

<222> 4

<223> Xaa = Ile, Leu or Val

<221> VARIANT

<222> 5

<223> Xaa = Asp, Asn or Glu

<221> VARIANT
 <222> 6,7
 <223> Xaa = any amino acid

<221> VARIANT
 <222> 8
 <223> Xaa = Ser, Thr or Asn

<400> 1311
 Xaa Phe Xaa Xaa Xaa Xaa Xaa Gly
 1 5

<210> 1312
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Representative claudin CAR sequence

<400> 1312
 Ile Tyr Ser Tyr
 1

<210> 1313
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Representative claudin CAR sequence

<400> 1313
 Thr Ser Ser Tyr
 1

<210> 1314
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Representative claudin CAR sequence

<400> 1314
 Val Thr Ala Phe
 1

<210> 1315
 <211> 4

<212> PRT
 <213> Artificial Sequence

<220>
 <223> Representative claudin CAR sequence

<400> 1315
 Val Ser Ala Phe
 1

<210> 1316
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1316
 Cys Gly Trp Val Met Asn Gln Gly Trp Val Met Asn Gln Cys
 1 5 10

<210> 1317
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1317
 Cys Arg Trp Ala Pro Ile Pro Arg Trp Ala Pro Ile Pro Cys
 1 5 10

<210> 1318
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1318
 Cys Gly Trp Val Met Asn Gln Gln Asn Met Val Trp Gly Cys
 1 5 10

<210> 1319

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1319

Cys Gln Asn Met Val Trp Gly Gly Trp Val Met Asn Gln Cys
 1 5 10

<210> 1320

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1320

Cys Arg Trp Ala Pro Ile Pro Pro Ile Pro Ala Trp Arg Cys
 1 5 10

<210> 1321

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Trp-containing CAR sequence in the cyclic peptides
 that may be linked in tandem.

<400> 1321

Cys Pro Ile Pro Ala Trp Arg Arg Trp Ala Pro Ile Pro Cys
 1 5 10

<210> 1322

<211> 5
 <212> PRT
 <213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<400> 1322

Cys Gly Trp Val Cys
 1 5

<210> 1323
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1323
Cys Gly Trp Val Trp Asn Gln Cys
1 5

<210> 1324
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1324
Cys Gly Trp Val Trp Asn Cys
1 5

<210> 1325
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1325
Cys Arg Gly Trp Val Cys
1 5

<210> 1326
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1326
Cys Arg Gly Trp Val Trp Cys
1 5

<210> 1327
<211> 6

<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1327
Cys Gly Trp Val Cys Asn
1 5

<210> 1328
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1328
Cys Gly Trp Val
1

<210> 1329
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1329
Cys Arg Gly Trp Val Trp Asn Gln Phe Cys
1 5 10

<210> 1330
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1330
Cys Arg Gly Trp Val Trp Asn Gln Phe Phe Cys
1 5 10

<210> 1331
<211> 10
<212> PRT
<213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<221> MOD_RES

<222> 2

<223> Xaa = beta,beta-tetramethylene cysteine

<400> 1331

Ile	Xaa	Gly	Trp	Val	Trp	Asn	Gln	Cys	Glu
1				5					10

<210> 1332

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<221> MOD_RES

<222> 2

<223> Xaa = beta,beta -pentamethylene cysteine

<400> 1332

Ile	Xaa	Gly	Trp	Val	Trp	Asn	Gln	Cys
1				5				

<210> 1333

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<400> 1333

Gly	Trp	Val	Trp	Asn	Gln	Pro	Cys
1				5			

<210> 1334

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<400> 1334

Cys	Arg	Trp	Ala	Pro	Cys
1				5	

<210> 1335
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1335
Cys Arg Trp Ala Pro Ile Pro Cys
1 5

<210> 1336
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1336
Cys Arg Trp Ala Pro Ile Cys
1 5

<210> 1337
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1337
Cys Arg Trp Ala Pro Ile Pro Cys Cys
1 5

<210> 1338
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1338
Cys Arg Trp Ala Pro Ile Pro Cys Ser Cys Met
1 5 10

<210> 1339
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1339
Cys Arg Trp Ala Cys Asn
1 5

<210> 1340
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<221> MOD_RES
<222> 5
<223> Xaa = penicillamine

<400> 1340
Cys Arg Trp Ala Xaa
1 5

<210> 1341
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1341
Cys Arg Trp Ala Pro Ile Pro Cys Ser Cys
1 5 10

<210> 1342
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1342
Cys Arg Trp Ala Pro Ile Pro Cys Ser Met Cys
1 5 10

<210> 1343
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<221> MOD_RES
<222> 2
<223> Xaa = beta,beta-tetramethylene cysteine

<400> 1343
Ile Xaa Arg Trp Ala Pro Ile Pro Cys Glu
1 5 10

<210> 1344
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<221> MOD_RES
<222> 2
<223> Xaa = beta,beta-pentamethylene cysteine

<400> 1344
Ile Xaa Arg Trp Ala Pro Ile Pro Cys
1 5

<210> 1345
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Peptide used in cyclization

<400> 1345
Arg Trp Ala Pro Ile Pro Cys Cys
1 5

<210> 1346
<211> 8
<212> PRT
<213> Artificial Sequence

<220>

<223> Peptide used in cyclization

<400> 1346

Lys Arg Trp Ala Pro Ile Pro Asp

1

5

<210> 1347

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide used in cyclization process

<400> 1347

Glu Asp Ala Cys

1

<210> 1348

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Peptide used in cyclization process

<400> 1348

Asp Cys Cys Ile

1

<210> 1349

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Modulating agent

<400> 1349

Ser His Ala Val Ser Ser

1

5

<210> 1350

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Modulating agent

<400> 1350

Ala His Ala Val Asp Ile
1 5

<210> 1351

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> N-cadherin CAR sequence

<400> 1351

Phe His Leu Arg Ala His Ala Val Asp Ile Asn Gly Asn Gln Val
1 5 10 15

<210> 1352

<211> 48

<212> PRT

<213> Artificial Sequence

<220>

<223> Occludin CAR sequence

<400> 1352

Gly Val Asn Pro Thr Ala Gln Ser Ser Gly Ser Leu Tyr Gly Ser Gln
1 5 10 15
Ile Tyr Ala Leu Cys Asn Gln Phe Tyr Thr Pro Ala Ala Thr Gly Leu
20 25 30
Tyr Val Asp Gln Tyr Leu Tyr His Tyr Cys Val Val Asp Pro Gln Glu
35 40 45

<210> 1353

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Trp-containing cell adhesion recognition sequence

<400> 1353

Gly Trp Val Trp Asn Gln
1 5

<210> 1354

<211> 6

<212> PRT

<213> Artificial Sequence

<220>
<223> Trp-containing cell adhesion recognition sequence

<400> 1354
Asp Trp Ile Trp Asn Gln
1 5

<210> 1355
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Trp-containing cell adhesion recognition sequence

<400> 1355
Ser Trp Met Trp Asn Gln
1 5

<210> 1356
<211> 4
<212> PRT
<213> qArtificial Sequence

<220>
<223> Trp-containing cell adhesion recognition sequence

<400> 1356
Trp Val Asn Gln
1

<210> 1357
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Trp-containing cell adhesion recognition sequence

<400> 1357
Gly Trp Met Trp Asn Gln
1 5

<210> 1358
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1358
Asp Val Asn Glu
1

<210> 1359
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1359
Asp Ile Asn Asp Asn
1 5

<210> 1360
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1360
Asp Val Asn Asp Asn
1 5

<210> 1361
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1361
Val Asp Phe Glu
1

<210> 1362
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1362

Asp Ala Asp Glu
1

<210> 1363
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1363
Asp Val Asp Glu
1

<210> 1364
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1364
Asp Glu Asn Asp Asn
1 5

<210> 1365
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1365
Asp Val Asn Asp Glu
1 5

<210> 1366
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1366
Leu Asn Tyr Glu
1

<210> 1367
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1367
Asp Gln Asn Asp Asn
1 5

<210> 1368
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1368
Asp Thr Asn Glu
1

<210> 1369
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1369
Glu Val Asn Glu
1

<210> 1370
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Calcium binding motif

<400> 1370
Asp Ile Asn Asp
1

<210> 1371
 <211> 110
 <212> PRT
 <213> unknown

<220>
 <223> Obcad sequence

<400> 1371
 Arg Ser Lys Arg Gly Trp Val Trp Asn Gln Phe Phe Val Ile Glu Glu
 1 5 10 15
 Tyr Thr Gly Pro Asp Pro Val Leu Val Gly Arg Leu His Ser Asp Ile
 20 25 30
 Asp Ser Gly Asp Gly Asn Ile Lys Tyr Ile Leu Ser Gly Glu Gly Ala
 35 40 45
 Gly Thr Ile Phe Val Ile Asp Asp Lys Ser Gly Asn Ile His Ala Thr
 50 55 60
 Lys Thr Leu Asp Arg Glu Glu Arg Ala Gln Tyr Thr Leu Met Ala Gln
 65 70 75 80
 Ala Val Asp Arg Asp Thr Asn Arg Pro Leu Glu Pro Pro Ser Glu Phe
 85 90 95
 Ile Val Lys Val Gln Asp Ile Asn Asp Asn Pro Pro Glu Phe
 100 105 110

<210> 1372
 <211> 108
 <212> PRT
 <213> Unknown

<220>
 <223> Cad5 sequence

<400> 1372
 Arg Gln Lys Arg Asp Trp Ile Trp Asn Gln Met His Ile Asp Glu Glu
 1 5 10 15
 Lys Asn Thr Ser Leu Pro His His Val Gly Lys Ile Lys Ser Ser Val
 20 25 30
 Ser Arg Lys Asn Ala Lys Tyr Leu Leu Lys Gly Glu Tyr Val Gly Lys
 35 40 45
 Val Phe Arg Val Asp Ala Glu Thr Gly Asp Val Phe Ala Ile Glu Arg
 50 55 60
 Leu Asp Arg Glu Asn Ile Ser Glu Tyr His Leu Thr Ala Val Ile Val
 65 70 75 80
 Asp Lys Asp Thr Gly Glu Asn Leu Glu Thr Pro Ser Ser Phe Thr Ile
 85 90 95
 Lys Val His Asp Val Asn Asp Asn Trp Pro Val Phe
 100 105

<210> 1373
 <211> 110
 <212> PRT
 <213> unknown

<220>

<223> Cad6 sequence

<400> 1373

Arg	Ser	Lys	Arg	Ser	Trp	Met	Trp	Asn	Gln	Phe	Phe	Leu	Leu	Glu	Glu
1				5					10					15	
Tyr	Thr	Gly	Ser	Asp	Tyr	Gln	Tyr	Val	Gly	Lys	Leu	His	Ser	Asp	Gln
			20					25					30		
Asp	Arg	Gly	Asp	Gly	Ser	Leu	Lys	Tyr	Ile	Leu	Ser	Gly	Asp	Gly	Ala
		35					40					45			
Gly	Asp	Leu	Phe	Ile	Ile	Asn	Glu	Asn	Thr	Gly	Asp	Ile	Gln	Ala	Thr
	50					55					60				
Lys	Arg	Leu	Asp	Arg	Glu	Glu	Lys	Pro	Val	Tyr	Ile	Leu	Arg	Ala	Gln
65					70				75						80
Ala	Ile	Asn	Arg	Arg	Thr	Gly	Arg	Pro	Val	Glu	Pro	Glu	Ser	Glu	Phe
			85					90						95	
Ile	Ile	Lys	Ile	His	Asp	Ile	Asn	Asp	Asn	Glu	Pro	Ile	Phe		
			100					105					110		

<210> 1374

<211> 110

<212> PRT

<213> unknown

<220>

<223> Cad7 sequence

<400> 1374

Arg	Thr	Lys	Arg	Ser	Trp	Val	Trp	Asn	Gln	Phe	Phe	Val	Leu	Glu	Glu
1				5					10					15	
Tyr	Met	Gly	Ser	Asp	Pro	Leu	Tyr	Val	Gly	Lys	Leu	His	Ser	Asp	Val
			20					25					30		
Asp	Lys	Gly	Asp	Gly	Ser	Ile	Lys	Tyr	Ile	Leu	Ser	Gly	Glu	Gly	Ala
		35					40					45			
Ser	Ser	Ile	Phe	Ile	Ile	Asp	Glu	Asn	Thr	Gly	Asp	Ile	His	Ala	Thr
	50					55					60				
Lys	Arg	Leu	Asp	Arg	Glu	Glu	Gln	Ala	Tyr	Tyr	Thr	Leu	Arg	Ala	Gln
65					70				75						80
Ala	His	Asp	Arg	Leu	Thr	Asn	Lys	Pro	Val	Glu	Pro	Glu	Ser	Glu	Phe
			85					90						95	
Val	Ile	Lys	Ile	Gln	Asp	Ile	Asn	Asp	Asn	Glu	Pro	Lys	Phe		
			100					105					110		

<210> 1375

<211> 110

<212> PRT

<213> unknown

<220>

<223> Cad8 sequence

<400> 1375

Arg	Ser	Lys	Arg	Gly	Trp	Val	Trp	Asn	Gln	Met	Phe	Val	Leu	Glu	Glu
1				5					10					15	
Phe	Ser	Gly	Pro	Glu	Pro	Ile	Leu	Val	Gly	Arg	Leu	His	Thr	Asp	Leu
			20					25					30		
Asp	Pro	Gly	Ser	Lys	Lys	Ile	Lys	Tyr	Ile	Leu	Ser	Gly	Asp	Gly	Ala
		35					40					45			
Gly	Thr	Ile	Phe	Gln	Ile	Asn	Asp	Val	Thr	Gly	Asp	Ile	His	Ala	Ile
	50					55				60					
Lys	Arg	Leu	Asp	Arg	Glu	Glu	Lys	Ala	Glu	Tyr	Thr	Leu	Thr	Ala	Gln
65					70				75						80
Ala	Val	Asp	Trp	Glu	Thr	Ser	Lys	Pro	Leu	Glu	Pro	Pro	Ser	Glu	Phe
				85					90					95	
Ile	Ile	Lys	Val	Gln	Asp	Ile	Asn	Asp	Asn	Ala	Pro	Glu	Phe		
			100					105					110		

<210> 1376

<211> 110

<212> PRT

<213> unknown

<220>

<223> Cad12 sequence

<400> 1376

Arg	Val	Lys	Arg	Gly	Trp	Val	Trp	Asn	Gln	Phe	Phe	Val	Leu	Glu	Glu
1				5					10					15	
Tyr	Val	Gly	Ser	Glu	Pro	Gln	Tyr	Val	Gly	Lys	Leu	His	Ser	Asp	Leu
			20					25					30		
Asp	Lys	Gly	Glu	Gly	Thr	Val	Lys	Tyr	Thr	Leu	Ser	Gly	Asp	Gly	Ala
		35					40					45			
Gly	Thr	Val	Phe	Thr	Ile	Asp	Glu	Thr	Thr	Gly	Asp	Ile	His	Ala	Ile
	50					55				60					
Arg	Ser	Leu	Asp	Arg	Glu	Glu	Lys	Pro	Phe	Tyr	Thr	Leu	Arg	Ala	Gln
65					70				75						80
Ala	Val	Asp	Ile	Glu	Thr	Arg	Lys	Pro	Leu	Glu	Pro	Glu	Ser	Glu	Phe
				85					90					95	
Ile	Ile	Lys	Val	Gln	Asp	Ile	Asn	Asp	Asn	Glu	Pro	Lys	Phe		
			100					105					110		

<210> 1377

<211> 110

<212> PRT

<213> unknown

<220>

<223> Cad14 sequence

<400> 1377

Arg	Pro	Lys	Arg	Gly	Trp	Val	Trp	Asn	Gln	Phe	Phe	Val	Leu	Glu	Glu
1				5					10					15	
His	Met	Gly	Pro	Asp	Pro	Gln	Tyr	Val	Gly	Lys	Leu	His	Ser	Asn	Ser

			20					25					30				
Asp	Lys	Gly	Asp	Gly	Ser	Val	Lys	Tyr	Ile	Leu	Thr	Gly	Glu	Gly	Ala		
		35					40					45					
Gly	Thr	Ile	Phe	Ile	Ile	Asp	Asp	Thr	Thr	Gly	Asp	Ile	His	Ser	Thr		
	50					55					60						
Lys	Ser	Leu	Asp	Arg	Glu	Gln	Lys	Thr	His	Tyr	Val	Leu	His	Ala	Gln		
65					70					75					80		
Ala	Ile	Asp	Arg	Arg	Thr	Asn	Lys	Pro	Leu	Glu	Pro	Glu	Ser	Glu	Phe		
			85					90						95			
Ile	Ile	Lys	Val	Gln	Asp	Ile	Asn	Asp	Asn	Ala	Pro	Lys	Phe				
			100					105					110				

<210> 1378

<211> 110

<212> PRT

<213> unknown

<220>

<223> PBcad sequence

<400> 1378

Arg	Val	Lys	Arg	Gly	Trp	Val	Trp	Asn	Gln	Phe	Phe	Val	Val	Glu	Glu		
1				5				10						15			
Tyr	Thr	Gly	Thr	Glu	Pro	Leu	Tyr	Val	Gly	Lys	Ile	His	Ser	Asp	Ser		
		20					25					30					
Asp	Glu	Gly	Asp	Gly	Thr	Ile	Lys	Tyr	Thr	Ile	Ser	Gly	Glu	Gly	Ala		
	35					40					45						
Gly	Thr	Ile	Phe	Leu	Ile	Asp	Glu	Leu	Thr	Gly	Asp	Ile	His	Ala	Thr		
	50				55					60							
Glu	Arg	Leu	Asp	Arg	Glu	Gln	Lys	Thr	Phe	Tyr	Thr	Leu	Arg	Ala	Gln		
65				70					75						80		
Ala	Arg	Asp	Arg	Ala	Thr	Asn	Arg	Leu	Leu	Glu	Pro	Glu	Ser	Glu	Phe		
			85					90						95			
Ile	Ile	Lys	Val	Gln	Asp	Ile	Asn	Asp	Ser	Glu	Pro	Arg	Phe				
			100					105					110				

<210> 1379

<211> 106

<212> PRT

<213> Homo sapiens

<400> 1379

Gly	Trp	Val	Trp	Asn	Gln	Phe	Phe	Val	Ile	Glu	Glu	Tyr	Thr	Gly	Pro		
1				5				10						15			
Asp	Pro	Val	Leu	Val	Gly	Arg	Leu	His	Ser	Asp	Ile	Asp	Ser	Gly	Asp		
		20					25					30					
Gly	Asn	Ile	Lys	Tyr	Ile	Leu	Ser	Gly	Glu	Gly	Ala	Gly	Thr	Ile	Phe		
	35					40					45						
Val	Ile	Asp	Asp	Lys	Ser	Gly	Asn	Ile	His	Ala	Thr	Lys	Thr	Leu	Asp		
	50				55					60							
Arg	Glu	Glu	Arg	Ala	Gln	Tyr	Thr	Leu	Met	Ala	Gln	Ala	Val	Asp	Arg		
65					70				75						80		

Asp Thr Asn Arg Pro Leu Glu Pro Pro Ser Glu Phe Ile Val Lys Val
 85 90 95
 Gln Asp Ile Asn Asp Asn Pro Pro Glu Phe
 100 105

<210> 1380
 <211> 106
 <212> PRT
 <213> Mus musculus

<400> 1380
 Gly Trp Val Trp Asn Gln Phe Phe Val Ile Glu Glu Tyr Thr Gly Pro
 1 5 10 15
 Asp Pro Val Leu Val Gly Arg Leu His Ser Asp Ile Asp Ser Gly Asp
 20 25 30
 Gly Asn Ile Lys Tyr Ile Leu Ser Gly Glu Gly Ala Gly Thr Ile Phe
 35 40 45
 Val Ile Asp Asp Lys Ser Gly Asn Ile His Ala Thr Lys Thr Leu Asp
 50 55 60
 Arg Glu Glu Arg Ala Gln Tyr Thr Leu Met Ala Gln Ala Val Asp Arg
 65 70 75 80
 Asp Thr Asn Arg Pro Leu Glu Pro Pro Ser Glu Phe Ile Val Lys Val
 85 90 95
 Gln Asp Ile Asn Asp Asn Pro Pro Glu Phe
 100 105

<210> 1381
 <211> 4
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Calcium binding motif

<400> 1381
 Val Asp Tyr Glu
 1

<210> 1382
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Calcium binding motif

<400> 1382
 Asp Asp Asn Asp Asn
 1 5

<210> 1383
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Calcium binding motif

<400> 1383
 Asp Tyr Asn Asp Asn
 1 5

<210> 1384
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Calcium binding motif

<400> 1384
 Asp Ser Asn Asp Asn
 1 5

<210> 1385
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 1385
 Glu Trp Ile Lys Phe Ala Ala Ala Cys Arg Glu Gly Glu Asp Asn Ser
 1 5 10 15
 Lys Arg Asn Pro Ile Ala Lys Ile His Ser Asp Cys Ala Ala Asn Gln
 20 25 30
 Gln Val Thr Tyr Arg Ile Ser Gly Val Gly Ile Asp Gln Pro Pro Tyr
 35 40 45
 Gly Ile Phe Val Ile Asn Gln Lys Thr Gly Glu Ile Asn Ile Thr Ser
 50 55 60
 Ile Val Asp Arg Glu Val Thr Pro Phe Phe Ile Ile Tyr Cys Arg Ala
 65 70 75 80
 Leu Asn Ser Met Gly Gln Asp Leu Glu Arg Pro Leu Glu Leu Arg Val
 85 90 95
 Arg Val Leu Asp Ile Asn Asp Asn Pro Pro Val Phe
 100 105

<210> 1386
 <211> 108
 <212> PRT
 <213> Bos tarus

<400> 1386

Glu Trp Ile Lys Phe Ala Ala Ala Cys Arg Glu Gly Glu Asp Asn Ser
 1 5 10 15
 Lys Arg Asn Pro Ile Ala Lys Ile His Ser Asp Cys Ala Ala Asn Gln
 20 25 30
 Gln Val Thr Tyr Arg Ile Ser Gly Val Gly Ile Asp Gln Pro Pro Tyr
 35 40 45
 Gly Ile Phe Val Ile Asn Gln Lys Thr Gly Glu Ile Asn Ile Thr Ser
 50 55 60
 Ile Val Asp Arg Glu Val Thr Pro Phe Phe Ile Ile Tyr Cys Arg Ala
 65 70 75 80
 Leu Asn Ser Leu Gly Gln Asp Leu Glu Lys Pro Leu Glu Leu Arg Val
 85 90 95
 Arg Val Leu Asp Ile Asn Asp Asn Pro Pro Val Phe
 100 105

<210> 1387
 <211> 110
 <212> PRT
 <213> Homo sapiens

<400> 1387
 Ala Trp Ile Thr Ala Pro Val Ala Leu Arg Glu Gly Glu Asp Leu Ser
 1 5 10 15
 Lys Lys Asn Pro Ile Ala Lys Ile His Ser Asp Leu Ala Glu Glu Arg
 20 25 30
 Gly Leu Lys Ile Thr Tyr Lys Tyr Thr Gly Lys Gly Ile Thr Glu Pro
 35 40 45
 Pro Phe Gly Ile Phe Val Phe Asn Lys Asp Thr Gly Glu Leu Asn Val
 50 55 60
 Thr Ser Ile Leu Asp Arg Glu Glu Thr Pro Phe Phe Leu Leu Thr Gly
 65 70 75 80
 Tyr Ala Leu Asp Ala Arg Gly Asn Asn Val Glu Lys Pro Leu Glu Leu
 85 90 95
 Arg Ile Lys Val Leu Asp Ile Asn Asp Asn Glu Pro Val Phe
 100 105 110

<210> 1388
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 1388
 Glu Trp Val Lys Phe Ala Lys Pro Cys Arg Glu Gly Glu Asp Asn Ser
 1 5 10 15
 Lys Arg Asn Pro Ile Ala Lys Ile Thr Ser Asp Tyr Gln Ala Thr Gln
 20 25 30
 Lys Ile Thr Tyr Arg Ile Ser Gly Val Gly Ile Asp Gln Pro Pro Phe
 35 40 45
 Gly Ile Phe Val Val Asp Lys Asn Thr Gly Asp Ile Asn Ile Thr Ala
 50 55 60
 Ile Val Asp Arg Glu Glu Thr Pro Ser Phe Leu Ile Thr Cys Arg Ala
 65 70 75 80

Leu Asn Ala Gln Gly Leu Asp Val Glu Lys Pro Leu Ile Leu Thr Val
 85 90 95
 Lys Ile Leu Asp Ile Asn Asp Asn Pro Pro Val Phe
 100 105

<210> 1389
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 1389
 Glu Trp Val Lys Phe Ala Lys Pro Cys Arg Glu Arg Glu Asp Asn Ser
 1 5 10 15
 Arg Arg Asn Pro Ile Ala Lys Ile Thr Ser Asp Phe Gln Lys Asn Gln
 20 25 30
 Lys Ile Thr Tyr Arg Ile Ser Gly Val Gly Ile Asp Gln Pro Pro Phe
 35 40 45
 Gly Ile Phe Val Val Asp Pro Asn Asn Gly Asp Ile Asn Ile Thr Ala
 50 55 60
 Ile Val Asp Arg Glu Glu Thr Pro Ser Phe Leu Ile Thr Cys Arg Ala
 65 70 75 80
 Leu Asn Ala Leu Gly Gln Asp Val Glu Arg Pro Leu Ile Leu Thr Val
 85 90 95
 Lys Ile Leu Asp Val Asn Asp Asn Pro Pro Ile Phe
 100 105

<210> 1390
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 1390
 Glu Trp Ile Lys Phe Ala Ala Ala Cys Arg Glu Gly Glu Asp Asn Ser
 1 5 10 15
 Lys Arg Asn Pro Ile Ala Lys Ile Arg Ser Asp Cys Glu Ser Asn Gln
 20 25 30
 Lys Ile Thr Tyr Arg Ile Ser Gly Val Gly Ile Asp Arg Pro Pro Tyr
 35 40 45
 Gly Val Phe Thr Ile Asn Pro Arg Thr Gly Glu Ile Asn Ile Thr Ser
 50 55 60
 Val Val Asp Arg Glu Ile Thr Pro Leu Phe Leu Ile Tyr Cys Arg Ala
 65 70 75 80
 Leu Asn Ser Arg Gly Glu Asp Leu Glu Arg Pro Leu Glu Leu Arg Val
 85 90 95
 Lys Val Met Asp Ile Asn Asp Asn Ala Pro Val Phe
 100 105

<210> 1391
 <211> 108
 <212> PRT
 <213> Mus musculus

<400> 1391

Glu	Trp	Ile	Lys	Phe	Ala	Ala	Ala	Cys	Arg	Glu	Gly	Glu	Asp	Asn	Ser
1				5				10						15	
Lys	Arg	Asn	Pro	Ile	Ala	Arg	Ile	Arg	Ser	Asp	Cys	Glu	Val	Ser	Gln
		20						25					30		
Arg	Ile	Thr	Tyr	Arg	Ile	Ser	Gly	Ala	Gly	Ile	Asp	Arg	Pro	Pro	Tyr
		35					40					45			
Gly	Val	Phe	Thr	Ile	Asn	Pro	Arg	Thr	Gly	Glu	Ile	Asn	Ile	Thr	Ser
	50					55					60				
Val	Val	Asp	Arg	Glu	Ile	Thr	Pro	Leu	Phe	Leu	Ile	His	Cys	Arg	Ala
65					70					75					80
Leu	Asn	Ser	Arg	Gly	Glu	Asp	Leu	Glu	Arg	Pro	Leu	Glu	Leu	Arg	Val
				85					90					95	
Lys	Val	Met	Asp	Val	Asn	Asp	Asn	Pro	Pro	Val	Phe				
			100					105							

<210> 1392

<211> 108

<212> PRT

<213> Mus musculus

<400> 1392

Glu	Trp	Ile	Lys	Phe	Ala	Ala	Ala	Cys	Arg	Glu	Gly	Glu	Asp	Asn	Ser
1				5				10						15	
Lys	Arg	Asn	Pro	Ile	Ala	Lys	Ile	His	Ser	Asp	Cys	Ala	Ala	Asn	Gln
		20						25					30		
Pro	Val	Thr	Tyr	Arg	Ile	Ser	Gly	Val	Gly	Ile	Asp	Gln	Pro	Pro	Tyr
		35					40					45			
Gly	Ile	Phe	Ile	Ile	Asn	Gln	Lys	Thr	Gly	Glu	Ile	Asn	Ile	Thr	Ser
	50					55					60				
Ile	Val	Asp	Arg	Glu	Val	Thr	Pro	Phe	Phe	Ile	Ile	Tyr	Cys	Arg	Ala
65					70					75					80
Leu	Asn	Ala	Gln	Gly	Gln	Asp	Leu	Glu	Asn	Pro	Leu	Glu	Leu	Arg	Val
				85					90					95	
Arg	Val	Met	Asp	Ile	Asn	Asp	Asn	Pro	Pro	Val	Phe				
			100					105							

<210> 1393

<211> 108

<212> PRT

<213> Mus musculus

<400> 1393

Glu	Trp	Ile	Lys	Phe	Ala	Ala	Ala	Cys	Arg	Glu	Gly	Glu	Asp	Asn	Ser
1				5				10						15	
Lys	Arg	Asn	Pro	Ile	Ala	Lys	Ile	His	Ser	Asp	Cys	Ala	Ala	Asn	Gln
		20						25					30		
Pro	Val	Thr	Tyr	Arg	Ile	Ser	Gly	Val	Gly	Ile	Asp	Gln	Pro	Pro	Tyr
		35					40					45			
Gly	Ile	Phe	Ile	Ile	Asn	Gln	Lys	Thr	Gly	Glu	Ile	Asn	Ile	Thr	Ser
	50					55					60				

```

Ile Val Asp Arg Glu Val Thr Pro Phe Phe Ile Ile Tyr Cys Arg Ala
65          70          75          80
Leu Asn Ala Gln Gly Gln Asp Leu Glu Asn Pro Leu Glu Leu Arg Val
      85          90          95
Arg Val Met Asp Ile Asn Asp Asn Pro Pro Val Phe
      100          105

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<210> 1394
<211> 108
<212> PRT
<213> Homo sapiens

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<400> 1394
Arg Trp Ala Pro Ile Pro Ala Ser Leu Met Glu Asn Ser Leu Gly Pro
 1          5          10          15
Phe Pro Gln His Val Gln Gln Ile Gln Ser Asp Ala Ala Gln Asn Tyr
      20          25          30
Thr Ile Phe Tyr Ser Ile Ser Gly Pro Gly Val Asp Lys Glu Pro Phe
      35          40          45
Asn Leu Phe Tyr Ile Glu Lys Asp Thr Gly Asp Ile Phe Cys Thr Arg
      50          55          60
Ser Ile Asp Arg Glu Lys Tyr Glu Gln Phe Ala Leu Tyr Gly Tyr Ala
65          70          75          80
Thr Thr Ala Asp Gly Tyr Ala Pro Glu Tyr Pro Leu Pro Leu Ile Ile
      85          90          95
Lys Ile Glu Asp Asp Asn Asp Asn Ala Pro Tyr Phe
      100          105

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```

<210> 1395
<211> 108
<212> PRT
<213> Mus musculus

```

```

<400> 1395
Arg Trp Ala Pro Ile Pro Cys Ser Leu Met Glu Asn Ser Leu Gly Pro
 1          5          10          15
Phe Pro Gln His Ile Gln Gln Ile Gln Ser Asp Ala Ala Gln Asn Tyr
      20          25          30
Thr Ile Phe Tyr Ser Ile Ser Gly Pro Gly Val Asp Lys Glu Pro Tyr
      35          40          45
Asn Leu Phe Tyr Ile Glu Lys Asp Thr Gly Asp Ile Tyr Cys Thr Arg
      50          55          60
Ser Ile Asp Arg Glu Gln Tyr Asp Gln Phe Leu Val Tyr Gly Tyr Ala
65          70          75          80
Thr Thr Ala Asp Gly Tyr Ala Pro Asp Tyr Pro Leu Pro Leu Leu Phe
      85          90          95
Lys Val Glu Asp Asp Asn Asp Asn Ala Pro Tyr Phe
      100          105

```

```

<210> 1396
<211> 108

```

<212> PRT
 <213> Bos tarus

<400> 1396

Arg	Trp	Ala	Pro	Ile	Pro	Cys	Ser	Leu	Met	Glu	Asn	Ser	Leu	Gly	Pro
1				5					10					15	
Phe	Pro	Gln	His	Val	Gln	Gln	Val	Gln	Ser	Asp	Ala	Ala	Gln	Asn	Tyr
		20						25					30		
Thr	Ile	Phe	Tyr	Ser	Ile	Ser	Gly	Pro	Gly	Val	Asp	Lys	Glu	Pro	Phe
	35						40					45			
Asn	Leu	Phe	Phe	Ile	Glu	Lys	Asp	Thr	Gly	Asp	Ile	Phe	Cys	Thr	Arg
	50					55					60				
Ser	Ile	Asp	Arg	Glu	Gln	Tyr	Gln	Glu	Phe	Pro	Ile	Tyr	Ala	Tyr	Ala
65					70					75					80
Thr	Thr	Ala	Asp	Gly	Tyr	Ala	Pro	Glu	Tyr	Pro	Leu	Pro	Leu	Val	Phe
			85						90					95	
Lys	Val	Glu	Asp	Asp	Asn	Asp	Asn	Ala	Pro	Tyr	Phe				
			100					105							

<210> 1397
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 1397

Arg	Trp	Ala	Pro	Ile	Pro	Cys	Ser	Met	Leu	Glu	Asn	Ser	Leu	Gly	Pro
1				5					10					15	
Phe	Pro	Leu	Phe	Leu	Gln	Gln	Val	Gln	Ser	Asp	Thr	Ala	Gln	Asn	Tyr
		20						25					30		
Thr	Ile	Tyr	Tyr	Ser	Ile	Arg	Gly	Pro	Gly	Val	Asp	Gln	Glu	Pro	Arg
	35						40					45			
Asn	Leu	Phe	Tyr	Val	Glu	Arg	Asp	Thr	Gly	Asn	Leu	Tyr	Cys	Thr	Arg
	50					55					60				
Pro	Val	Asp	Arg	Glu	Gln	Tyr	Glu	Ser	Phe	Glu	Ile	Ile	Ala	Phe	Ala
65					70					75					80
Thr	Thr	Pro	Asp	Gly	Tyr	Thr	Pro	Glu	Leu	Pro	Leu	Pro	Leu	Ile	Ile
			85						90					95	
Lys	Ile	Glu	Asp	Glu	Asn	Asp	Asn	Tyr	Pro	Ile	Phe				
			100					105							

<210> 1398
 <211> 108
 <212> PRT
 <213> Canis familiaris

<400> 1398

Arg	Trp	Ala	Pro	Ile	Pro	Cys	Ser	Met	Gln	Glu	Asn	Ser	Leu	Gly	Pro
1				5					10					15	
Phe	Pro	Leu	Phe	Leu	Gln	Gln	Ile	Gln	Ser	Asp	Thr	Ala	Gln	Asn	Tyr
		20						25					30		
Thr	Ile	Phe	Tyr	Ser	Ile	Arg	Gly	Pro	Gly	Val	Asp	Arg	Glu	Pro	Lys
	35						40					45			

```

Asn Leu Phe Tyr Val Glu Arg Asp Thr Gly Asn Leu Phe Cys Thr Arg
 50          55          60
Pro Val Asp Arg Glu Glu Tyr Glu Ser Phe Glu Leu Ile Ala Phe Ala
65          70          75          80
Thr Thr Pro Asp Gly Tyr Thr Pro Glu Leu Pro Leu Pro Leu Val Ile
          85          90          95
Arg Ile Glu Asp Glu Asn Asp Asn Tyr Pro Ile Phe
          100          105

```

<210> 1399
 <211> 108
 <212> PRT
 <213> Homo sapiens

```

<400> 1399
Arg Trp Ala Pro Ile Pro Cys Ser Met Gln Glu Asn Ser Leu Gly Pro
 1          5          10          15
Phe Pro Leu Phe Leu Gln Gln Val Glu Ser Asp Ala Ala Gln Asn Tyr
          20          25          30
Thr Val Phe Tyr Ser Ile Ser Gly Arg Gly Val Asp Lys Glu Pro Leu
          35          40          45
Asn Leu Phe Tyr Ile Glu Arg Asp Thr Gly Asn Leu Phe Cys Thr Arg
          50          55          60
Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
65          70          75          80
Ser Thr Ala Asp Gly Tyr Ser Ala Asp Leu Pro Leu Pro Leu Pro Ile
          85          90          95
Arg Val Glu Asp Glu Asn Asp Asn His Pro Val Phe
          100          105

```

<210> 1400
 <211> 108
 <212> PRT
 <213> Mus musculus

```

<400> 1400
Arg Trp Ala Pro Ile Pro Cys Ser Met Gln Glu Asn Ser Leu Gly Pro
 1          5          10          15
Phe Pro Leu Phe Leu Gln Gln Val Gln Ser Asp Ala Ala Gln Asn Tyr
          20          25          30
Thr Val Phe Tyr Ser Ile Ser Gly Arg Gly Ala Asp Gln Glu Pro Leu
          35          40          45
Asn Trp Phe Phe Ile Glu Arg Asp Thr Gly Asn Leu Tyr Cys Thr Arg
          50          55          60
Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
65          70          75          80
Ser Thr Ala Asp Gly Tyr Ser Ala Asp Leu Pro Leu Pro Leu Pro Ile
          85          90          95
Lys Ile Glu Asp Glu Asn Asp Asn Tyr Pro Leu Phe
          100          105

```


<210> 1401
 <211> 108
 <212> PRT
 <213> Bos tarus

<400> 1401
 Arg Trp Ala Pro Ile Pro Cys Ser Met Gln Glu Asn Ser Leu Gly Pro
 1 5 10 15
 Phe Pro Leu Phe Leu Gln Gln Val Gln Ser Asp Ala Ala Gln Asn Tyr
 20 25 30
 Thr Ile Phe Tyr Ser Ile Ser Gly Arg Gly Val Asp Lys Glu Pro Leu
 35 40 45
 Asn Leu Phe Phe Ile Glu Arg Asp Thr Gly Asn Leu Tyr Cys Thr Gln
 50 55 60
 Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
 65 70 75 80
 Ser Thr Ala Asp Gly Tyr Ser Ala Asp Phe Pro Leu Pro Leu Pro Ile
 85 90 95
 Arg Val Glu Asp Glu Asn Asp Asn His Pro Ile Phe
 100 105

<210> 1402
 <211> 108
 <212> PRT
 <213> Homo sapiens

<400> 1402
 Arg Trp Ala Pro Ile Pro Cys Ser Met Gln Glu Asn Ser Leu Gly Pro
 1 5 10 15
 Phe Pro Leu Phe Leu Gln Gln Val Glu Ser Asp Ala Ala Gln Asn Tyr
 20 25 30
 Thr Val Phe Tyr Ser Ile Ser Gly Arg Gly Val Asp Lys Glu Pro Leu
 35 40 45
 Asn Leu Phe Tyr Ile Glu Arg Asp Thr Gly Asn Leu Phe Cys Thr Arg
 50 55 60
 Pro Val Asp Arg Glu Glu Tyr Asp Val Phe Asp Leu Ile Ala Tyr Ala
 65 70 75 80
 Ser Thr Ala Asp Gly Tyr Ser Ala Asp Leu Pro Leu Pro Leu Pro Ile
 85 90 95
 Arg Val Glu Asp Glu Asn Asp Asn His Pro Val Phe
 100 105